

Amendments to the Claims:

This listing of claims replaces all prior versions and listings of claims in the application:

Listing of Claims:

1.-57. (Cancelled).

58. (Currently amended) An isolated polynucleotide comprising a nucleic acid encoding a polypeptide having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO:2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2; and

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2,

said polypeptide effective for catalysing the hydroxylation of campestanol.

59. (Previously presented) The polynucleotide of claim 58, wherein said polypeptide has from 85% to 90% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

60. (Previously presented) The polynucleotide of claim 58, wherein said polypeptide has from 90% to 95% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

61. (Previously presented) The polynucleotide of claim 58, wherein said polypeptide has from 95% to 98% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

62. (Previously presented) The polynucleotide of claim 58, wherein said polypeptide has the amino acid sequence set forth in SEQ ID NO: 2.

63. (Previously presented) The polynucleotide of claim 58, wherein said polynucleotide further comprises a control element operably linked to said nucleic acid encoding said polypeptide.

64. (Previously presented) The polynucleotide of claim 63, wherein said control element is a tissue-specific promoter.

65. (Previously presented) The polynucleotide of claim 63, wherein said control element is an embryonic storage protein promoter.

66. (Previously presented) The polynucleotide of claim 63, wherein said control element comprises nucleotides 2102 to 3202 of SEQ ID NO: 1.

67. (Currently amended) An isolated polynucleotide comprising a nucleic acid encoding a polypeptide having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2;

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2.

said polypeptide effective for catalysing the hydroxylation of 6-oxocampestanol.

68. (Previously presented) The polynucleotide of claim 67, wherein said polypeptide has from 85% to 90% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

69. (Previously presented) The polynucleotide of claim 67, wherein said polypeptide has from 90% to 95% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

70. (Previously presented) The polynucleotide of claim 67, wherein said polypeptide has from 95% to 98% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

71. (Previously presented) The polynucleotide of claim 67, wherein said polypeptide has the amino acid sequence set forth in SEQ ID NO: 2.

72. (Previously presented) The polynucleotide of claim 67, wherein said polynucleotide further comprises a control element operably linked to said nucleic acid encoding said polypeptide.

73. (Previously presented) The polynucleotide of claim 72, wherein said control element is a tissue-specific promoter.

74. (Previously presented) The polynucleotide of claim 72, wherein said control element is an embryonic storage protein promoter.

75. (Previously presented) The polynucleotide of claim 72, wherein said control element comprises nucleotides 2102 to 3202 of SEQ ID NO: 1.

76. (Currently amended) A transgenic plant containing at least one exogenous polynucleotide, said at least one exogenous polynucleotide comprising a nucleic acid encoding a polypeptide having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2;

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2, and

a control element operably linked to said nucleic acid encoding said polypeptide, wherein said polypeptide is effective for catalysing the hydroxylation of campestanol.

77. (Previously presented) The plant of claim 76, wherein said polypeptide has from 85% to 90% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

78. (Previously presented) The plant of claim 76, wherein said polypeptide has from 90% to 95% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

79. (Previously presented) The plant of claim 76, wherein said polypeptide has from 95% to 98% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

80. (Previously presented) The plant of claim 76, wherein said polypeptide has the amino acid sequence set forth in SEQ ID NO: 2.

81. (Cancelled).

82. (Currently amended) The plant of claim ~~81~~ 76, wherein said control element is a tissue-specific promoter.

83. (Currently amended) The plant of claim ~~81~~ 76, wherein said control element is an embryonic storage protein promoter.

84. (Currently amended) The plant of claim ~~81~~ 76, wherein said control element comprises nucleotides 2102 to 3202 of SEQ ID NO: 1.

85. (Currently amended) A transgenic plant containing at least one exogenous polynucleotide, said at least one exogenous polynucleotide comprising a nucleic acid encoding a polypeptide having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2; and

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2,

and wherein said polypeptide is effective for catalysing the hydroxylation of 6-oxocampestanol.

86. (Currently amended) A method of making a transgenic plant comprising introducing into a plant a polynucleotide comprising a nucleotide sequence encoding a polypeptide effective for catalysing the hydroxylation of campestanol and having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2; and

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2, thereby making said transgenic plant.

87. (Currently amended) A method of making a transgenic plant comprising introducing into a plant a polynucleotide comprising a nucleotide sequence encoding a polypeptide effective for catalysing the hydroxylation of 6-oxocampestanol and having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2;

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2, thereby making said transgenic plant.

88. (Previously presented) An isolated polynucleotide comprising nucleotides 2102 to 3202 of SEQ ID NO: 1.

89. (Previously presented) The isolated polynucleotide of claim 88, wherein said polynucleotide comprises nucleotides 1 to 3202 of SEQ ID NO: 1.

90. (Cancelled).

91. (Currently amended) A plant or bacterial host cell comprising the ~~exogenous~~ polynucleotide of claim 63.

92. (Currently amended) A plant or bacterial host cell comprising the ~~exogenous~~ polynucleotide of claim 72.
93. (Previously presented) A method of producing a polypeptide comprising the steps of:
(a) providing the host cell of claim 91; and
(b) culturing said host cell under conditions whereby said polypeptide encoded by said nucleic acid is expressed.
94. (Previously presented) A method of producing a polypeptide comprising the steps of:
(a) providing the host cell of claim 92; and
(b) culturing said host cell under conditions whereby said polypeptide encoded by said nucleic acid is expressed.
95. (New) The polynucleotide of claim 58, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.
96. (New) The polynucleotide of claim 58, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.
97. (New) The polynucleotide of claim 63, wherein said control element directs expression in vegetative tissue of a plant.
98. (New) The polynucleotide of claim 97, wherein said vegetative tissue is root tissue.
99. (New) The polynucleotide of claim 97, wherein said vegetative tissue is shoot tissue.
100. (New) The polynucleotide of claim 97, wherein said vegetative tissue is leaf tissue.

101. (New) The polynucleotide of claim 67, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

102. (New) The polynucleotide of claim 67, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

103. (New) The polynucleotide of claim 72, wherein said control element directs expression in vegetative tissue of a plant.

104. (New) The polynucleotide of claim 103, wherein said vegetative tissue is root tissue.

105. (New) The polynucleotide of claim 103, wherein said vegetative tissue is shoot tissue.

106. (New) The polynucleotide of claim 103, wherein said vegetative tissue is leaf tissue.

107. (New) The transgenic plant of claim 76, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

108. (New) The transgenic plant of claim 76, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

109. (New) The transgenic plant of claim 76, wherein said control element directs expression in vegetative tissue of a plant.

110. (New) The transgenic plant of claim 109, wherein said vegetative tissue is root tissue.

111. (New) The transgenic plant of claim 109, wherein said vegetative tissue is shoot tissue.

112. (New) The transgenic plant of claim 109, wherein said vegetative tissue is leaf tissue.

113. (New) The transgenic plant of claim 85, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

114. (New) The transgenic plant of claim 85, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

115. (New) The transgenic plant of claim 85, wherein said control element directs expression in vegetative tissue of a plant.

116. (New) The transgenic plant of claim 115, wherein said vegetative tissue is root tissue.

117. (New) The transgenic plant of claim 115, wherein said vegetative tissue is shoot tissue.

118. (New) The transgenic plant of claim 115, wherein said vegetative tissue is leaf tissue.

119. (New) The method of claim 86, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

120. (New) The method of claim 86, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

121. (New) The method of claim 87, wherein said polypeptide has from 55% to 60% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

122. (New) The method of claim 87, wherein said polypeptide has from 70% to 85% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2.

123. (New) An isolated polynucleotide comprising a nucleic acid encoding a polypeptide having

(a) greater than 43% sequence identity to the amino acid sequence set forth in SEQ ID NO: 2;

(b) 60% or greater sequence identity to domain A of SEQ ID NO:2; and

(c) 60% or greater sequence identity to domain B of SEQ ID NO:2,

said polypeptide effective for modulating one or more phenotypic traits selected from the group consisting of brassinosteroid synthesis, cell length, period of flowering, seed production, seed size, leaf size, and hypocotyl length.

124. (New) The isolated polynucleotide of claim 123, wherein said modulation is a decrease in one or more of said phenotypic traits.

125. (New) The isolated polynucleotide of claim 124, wherein said modulation is an increase in one or more of said phenotypic traits.

126. (New) The isolated polynucleotide of claim 123, wherein said phenotypic trait is a *dwf4* phenotypic trait.

127. (New) The isolated polynucleotide of claim 58 or 67, wherein said polypeptide has from 70% to 85% sequence identity to domain A of SEQ ID NO:2 and from 70% to 85% sequence identity to domain B of SEQ ID NO:2.

128. (New) The isolated polynucleotide of claim 127, wherein said polypeptide has from 85% to 90% sequence identity to domain A of SEQ ID NO:2 and from 85% to 90% sequence identity to domain B of SEQ ID NO:2.

129. (New) The isolated polynucleotide of claim 127, wherein said polypeptide has from 90% to 95% sequence identity to domain A of SEQ ID NO:2 and from 90% to 95% sequence identity to domain B of SEQ ID NO:2.

130. (New) The transgenic plant of claim 76 or 85, wherein said polypeptide has from 70% to 85% sequence identity to domain A of SEQ ID NO:2 and from 70% to 85% sequence identity to domain B of SEQ ID NO:2.

131. (New) The transgenic plant of claim 130, wherein said polypeptide has from 85% to 90% sequence identity to domain A of SEQ ID NO:2 and from 85% to 90% sequence identity to domain B of SEQ ID NO:2.

132. (New) The transgenic plant of claim 130, wherein said polypeptide has from 90% to 95% sequence identity to domain A of SEQ ID NO:2 and from 90% to 95% sequence identity to domain B of SEQ ID NO:2.